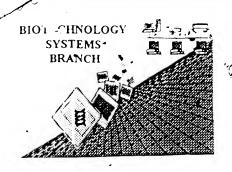
## RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09614221
Source:	OIPE
Date Processed by STIC:	11/16/01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 c-mail help: <a href="mailto:patin21help@uspto.gov">patin21help@uspto.gov</a> or phone 703-306-4119 (R. Wax) PATENTIN 3.0 c-mail help: <a href="mailto:patin3help@uspto.gov">patin3help@uspto.gov</a> or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## Checker Version 3.0

The Checker Version 3 0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3 0 works for sequence listings generated for the original version of 37 CFR §§1 821 – 1 825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

## Raw Sequence Litting Error Summary

ERROR DETECTED	SUCCESTED CORRECTION SERIAL NUMBER: 19/6/4 22/
ATTN: NEW RULES CAS	ES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY P
1Wrapped Nucleics Wrapped Aminos	The numberNext at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to 3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
)Hisaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use apace characters, instead.
1] ZA-non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
SVariable Length.	Sequence(s)contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable kingth and indicate in the <220>-<223> section that some may be missing.
6Patentin 2.0 "bug"	A "bug" in Patentin version 2.0 has caused the <220> <223> section to be missing from amino acid sequences(s).  Normally, Patentin would automatically, generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220> <223> section to the subsequent amino acid sequence. This applies to the mandatory <220> <223> sections for Artificial or Unknown sequence.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
, · · · · .	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to Include the skipped sequences
Skipped Sequences (NEW RULES)	Sequence(s) missing. If Intentional, please insert the following lines for each skipped sequen <210> sequence id number <400> sequence id number 000
9 Vuic of n'i or Xii'i (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing.  Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents
10Invalid <213> Response	Per 1.823 of Sequence Rules, the only valld <213> responses are: Unknown, Artificial Sequence, or scientific'name (Genustspecies) <220>-<223> section is required when <213> response is Unknown is Artificial Sequence
IUx of <270>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (See. 1.823 of Sequence Rules)
2Patentin 2.0 "bug"	Please do not use "Copy to Disk" function of Patentin version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
3 Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represe any value not specifically a nucleotide.

AMC/MH - Biotechnology Systems Branch - 08/21/2001

OIPE

RAW SEQUENCE LISTING DATE: 11/16/2001 PATENT APPLICATION: US/09/614,221 TIME: 10:41:23

Input Set : N:\jumbos\614221.raw

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2 <110> APPLICANT: Karunanandaa, Balasulojini
             Yu, Jaehyuk
             Kishore, Ganesh M.
      6 <120> TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED
             WITH STEROL SYNTHESIS AND METABOLISM
      9 <130> FILE REFERENCE: 05686.0004.NPUS00
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C--> 11 <141> CURRENT FILING DATE: 2000-07-11
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    12 <151> PRIOR FILING DATE: 1999-07-12
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    29
                                                                            101
    31 atg gag tac tot tac otg tta gat atg gog gac aag act gag gat coa
    32 Met Glu Tyr Ser Tyr Leu Leu Asp Met Ala Asp Lys Thr Glu Asp Pro
                                            20
    35 tac atg aga cta gta tat get tea tea tte ttt ata tet gte tae tat
                                                                            149
    36 Tyr Met Arg Leu Val Tyr Ala Ser Ser Phe Phe Ile Ser Val Tyr Tyr
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                    30
                                        35
    39 qcc tat caa cga acq tqq aaq cca ttc aat cca att ctt ggt gag act
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    40 Ala Tyr Gln Arg Thr Trp Lys Pro Phe Asn Pro Ile Leu Gly Glu Thr
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    44 Tyr Glu Met Val Asn His Gly Gly Ile Thr Phe Ile Ser Glu Gln Val
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                                                                            293
    47 agt cat cac cct cca atg agt gct ggg cat gct gaa act gaa cat ttc
    48 Ser His His Pro Pro Met Ser Ala Gly His Ala Glu Thr Glu His Phe
    49 75
                           80
                                                85
                                                                            341
    51 act tat gat gtt aca tca aaa ttg aaa acc aaa ttt ctc ggc aac tca
    52 Thr Tyr Asp Val Thr Ser Lys Leu Lys Thr Lys Phe Leu Gly Asn Ser
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                       95
                                                                            389
    55 gtt gat gta tat cct gtt gga aga acg cgt gtt acc ctc aaa aga gat
    56 Val Asp Val Tyr Pro Val Gly Arg Thr Arg Val Thr Leu Lys Arg Asp
                                                                            437
    59 ggt gtg gtc ctt gat ttg gtg cct cct cct aca aaa gtt agc aac ttg
    60 Gly Val Val Leu Asp Leu Val Pro Pro Pro Thr Lys Val Ser Asn Leu
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                                                                            485
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RAW SEQUENCE LISTING DATE: 11/16/2001 PATENT APPLICATION: US/09/614,221 TIME: 10:41:23

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69 155 160 165 170	
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72 Trp Phe Gly Tyr Glu Val Asp Gly Tyr Val Tyr Asn Ser Ala Asp Glu	
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75 cct aag ata ctg atg act gga aaa tgg aat gag gct atg aat tat caa	629
76 Pro Lys Ile Leu Met Thr Gly Lys Trp Asn Glu Ala Met Asn Tyr Gln	
77 190 195 200	
79 gtt tgt gac tca gag gga gaa cca ctt cca ggc act gag ttg aaa gag	677
80 Val Cys Asp Ser Glu Gly Glu Pro Leu Pro Gly Thr Glu Leu Lys Glu	
81 205 210 215	705
83 att tgg aga gtt gct gat acc ccg aag aag gac aag ttc cag tac acg	725
84 Ile Trp Arg Val Ala Asp Thr Pro Lys Lys Asp Lys Phe Gln Tyr Thr 85 220 225 230	
85 220 230 87 cat ttt gca cac aag att aac agc ttt gac act gct ccc aag aag ttg	773
88 His Phe Ala His Lys Ile Asn Ser Phe Asp Thr Ala Pro Lys Lys Leu	775
89 235 240 245 250	
91 ttg gca tct gac tct cgt cta cgt cct gat aga atg gcc ctt gag aag	821
92 Leu Ala Ser Asp Ser Arg Leu Arg Pro Asp Arg Met Ala Leu Glu Lys	
93 255 260 265	
95 ggt gac cta tcc aca tct ggt tat gag aag agc agt ttg gag gag agg	869
96 Gly Asp Leu Ser Thr Ser Gly Tyr Glu Lys Ser Ser Leu Glu Glu Arg	
97 270 275 280	
99 caa aga gct gag aag aga aac cga gag gcc aag ggc cat aag ttc act	917
100 Gln Arg Ala Glu Lys Arg Asn Arg Glu Ala Lys Gly His Lys Phe Thr	
101 285 290 295	
103 cct aga tgg ttt gat tta aca gat gaa gta act cct acc cct tgg ggt	965
104 Pro Arg Trp Phe Asp Leu Thr Asp Glu Val Thr Pro Trp Gly	
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10% gae try gaa get tae daa tae aac ggt daa tat acc daa dat tgt get 108 Asp Leu Glu Val Tyr Gln Tyr Asn Gly Lys Tyr Thr Gln His Cys Ala	1013
109 315 320 325 330	
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/614,221

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143															111		
144				1	ec C	ys n.	on As	511 G. 5	Ly G.	111 50	J. F.	LO DO	10		rg ri	ic iic	
	tot	ata	αta	_	taa	tac	ata	tet	acc	act	cac	cct		_	+++	aat	159
							Ile										
149		15					20					25					007
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							ctt										447
		мет	Asn	Cys	Pro	_	Leu	ьeu	Leu	Arg		Leu	Pro	Val	PIO		
	110					115				~~~	120	-+-	~~~		~~+	125	495
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	Ala	ASP	ттр	Ald	130	1111	Val	ASII	TTG	135	Cys	Leu	GIU	1111	140	Leu	
177	a+ >	aat	~~~	++-		+ 2.0	aga	+ 02	a a t		+++	ot a	aa.	a++	•	aaa	543
							Arg										343
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RAW SEQUENCE LISTING

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DATE: 11/16/2001

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203 agc aaa gac tgg gag aaa gca aga gaa gca agc cta aaa gtt gag gaa   831		GIN	Inr	GIU		Ala	HIS	val	тгр		GIU	Leu	ASII	GIII		rre	vai	
204 Ser Lys Asp Trp Glu Lys Ala Arg Glu Ala Lys Leu Lys Val Glu Glu 205 240 245 250 250 260 250 260 265 260 265 260 265 260 265 265 260 265 265 260 265 265 265 260 265 265 265 265 265 265 265 265 265 265	-: 0.3			~~~		~~~	222	~~~	202		~~~	224	at a	222		a a a	~ a a	Q 3 1
205	204	age	Tuo	yac	Trn	Clu	Tuc	λla	Ara	Clu	γCa	Tuc	Lou	Tye	Val	Glu	Glu	031
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208 Arg Gln Arg Glu Leu Val Arg Glu Arg Glu Ser Lys Gly Glu Thr Trp 209											~~~	+ ~ ~	222		~ · · ·	202	+ ~ ~	970
209																		0/9
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212 Ile Ser Lys His Phe Val Val Ser Asn Asn Lys Glu Gly Trp Gln Cys 213 270 275 280 285 215 tca cct att cat aag agt gta cct gcg gcc cc atc aca ag gcc cta taa 285 216 Ser Pro Ile His Lys Ser Val Pro Ala Ala Pro Ile Thr Ala Leu 217 290 295 300 219 ttgttgtcac tgtcaagtag tgtaaagcat taaagtacat tttagaagag aatgttcata 210 aaaaattta atggttgaaa ttttgacaac aatgaagtact ataaacaaaat ttaaaattag 211 aaaaaattta atggttgaaa ttttgacaac aatgaagtact ataaacaaaat ttaaaaattag 212 aaaaaattta aagatgaaa aaaaaaaaag ggcggccgc g 213 ttacaatttt aaaaaaaaaa aaaaaaaaag ggcggccgc g 214 tcaatttt aaaaaaaaaa aaaaaaaaag ggcggccgc g 215 tea Colo SeQ ID NO: 3 227 <211> LENGTH: 1355 228 <212> TYPE: DNA 229 <213> ORGANISM: Glycine max 231 <220> FEATURE: 232 <221> NAME/KEY: CDS 233 <222> LOCATION: (32)(1099) 235 <400> SEQUENCE: 3 237 ggaattcggc tcgaggacaa tgcttcagaa a atg gct gag ctt atg gag tac 238 Met Ala Glu Leu Met Glu Tyr 239 1 5 241 tct tac ctg tta gat atg gcg gac aag aag act gag gat cca tac atg aga 242 Ser Tyr Leu Leu Asp Met Ala Asp Lys Thr Glu Asp Pro Tyr Met Arg 243 10 15 245 cta gta tat gct tca tca ttc ttt ata tct gtc tac tat gcc tat caa 246 Leu Val Tyr Ala Ser Ser Phe Phe Ile Ser Val Tyr Tyr Ala Tyr Gln 247 25 30 249 cga acg tgg aag cca ttc aat cca att ctt ggt gag act tat gaa atg 250 Arg Thr Trp Lys Pro Phe Asn Pro Ile Leu Gly Glu Thr Tyr Glu Met 251 40 45 50 253 gtt aac cat ggt ggc att aca ttt ata tca gag cag gtc agt cat cac 244 254 Val Asn His Gly Gly Ile Thr Phe Ile Ser Glu Gln Val Ser His His 255 60 70 257 cct cca atg agt gct ggg cat gct gaa act ga cat tca cat tat gat 258 Pro Pro Met Ser Ala Gly His Ala Glu Thr Glu His Phe Thr Tyr Asp 256 19tt aca tca act ga gtg aac caa ttc caa ttc ctc gcc act cac tac cac 258 Pro Pro Met Ser Ala Gly His Ala Glu Thr Glu His Phe Thr Tyr Asp 256 19tt aca tca act caa act ggt gaa act tca aca ttc ctc tcc cac act gct gas act cac act															+		+ ~+	027
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289	gtt	tgt	gac	tca	gag	gga	gaa	cca	ctt	cca	ggc	act	gag	ttg	aaa	gag	676
				Ser													
291		-	-			205					210					215	
		taa	aga	gtt	qct	gat	acc	ccq	aaq	aaq	gac	aag	ttc	caq	tac	acg	724
				Val													
295		-	,		220	•			•	225	-	-			230		
	cat.	ttt	gca	cac	aaq	att	aac	agc	ttt	gac	act	qct	ccc	aaq	aaq	ttq	772
				His													
299				235	-1-				240					245	-		
	tta	gca	tct		t.ct.	cat	cta	cat		gat	aσa	atσ	acc	ctt	gag	aag.	820
				Asp													
303	204		250			9		255			5		260			2	
	aat	gac		tcc	aca	tet	aat.		σασ	aaσ	age	agt	tta	gag	gag	agg	868
				Ser													
307	J_1	265					270	- 1 -		1		275				,	
	caa		act.	gag	aaσ	aga		cσa	σασ	acc	aaσ	aac	cat	aaσ	ttc	act	916
				Glu													
311		111 9	1114	014	210	285		9	014		290	0.1		-15	•	295	
		ада	taa	ttt	πat		aca	crat	gaa	αta		cct	acc	aat	taa		964
				Phe													
315	110	AIG	111	riic	300	пси	1111	пор	Olu	305	1111	110	1111		310		
	α <b>2</b> Ω	++~	m = =	gtt		caa	tac	220	aat		tat	acc	caa	cat		act	1012
				Val													1012
319	кър	ьеи	GIU	315	1 1 1	GIII	1 7 1	ASII	320	цуз	1 Y 1	1111	GIII	325	Cys	AIU	
	aaa	~++	a +	agt	+ a+	~ ~ ~	+ 00	a++		ata	act	a a c	ato		cca	αaa	1060
				Ser													1000
323	нта	val	330	ser	261	Gru	Cys	335	Giu	val	FIU	дор	340	Arg	110	Gia	
	++ ~	220		tgg		+ = +	ant.		++~	ant.	aat	<i>α</i> 2 2		tasa	roato	act	1109
													Lay	cya	<i>y</i> ca co		1109
326	PHe		PIO	Trp	GIII	1 7 1		ASII	Leu	ASP	нта	355					
327		345					350		4				. ~ + + +			+	1160
		_					-									taatct	1169
																atttga	1229
																agtaga	1289
335	gaaa	attaa	aag t	taaa	aaaa	aa aa	aaaa	aaaa	a aaa	aaaa	aaaa	aaaa	aaaaa	aaa a	aaaa	gggcgg	1349

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/614,221 TIME: 10:41:24

DATE: 11/16/2001

Input Set : N:\jumbos\614221.raw

```
L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:644 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:7
L:644 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:7
L:644 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:646 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:7
L:646\ M:258\ W: Mandatory Feature missing, <222> not found for SEQ ID#:7
L:646 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:648 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:7
L:648\ M:258\ W: Mandatory Feature missing, <222> not found for SEQ ID#:7
\text{L:}648\ \text{M:}341\ \text{W:} (46) "n" or "Xaa" used, for SEQ ID#:7
L:661 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:8
L:661\ M:258\ W: Mandatory Feature missing, <222> not found for SEQ ID#:8
L:661 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:663 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:8
L:663 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:8
L:663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:665 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:8
L:665\ M:258\ W: Mandatory Feature missing, <222> not found for SEQ ID#:8
L:665 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:667 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:8
L:667 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:8
L:667 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:718 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:11
L:718\ M:258\ W: Mandatory Feature missing, <222> not found for SEQ ID#:11
L:718 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:757 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:13
L:757 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:757 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:763 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:13
L:763 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:763 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:778 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:14
L:778 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:778 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:788 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:14
L:788 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:788\ M:341\ W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:809 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:15
L:809 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:15
L:809 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:844 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:17
L:844 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:17
L:844 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:846 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:17
L:846 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:17
L:846 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:923 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:21
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DATE: 11/16/2001

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/614,221 TIME: 10:41:24

Input Set : N:\jumbos\614221.raw

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L:923 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
L:923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:992 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:24
L:992 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
L:992 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:994 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:24
L:994 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
L:994 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:996 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:24
L:996 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
L:996 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:998 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:24
L:998 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
L:998 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:1013 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:1013 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
L:1013 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1015 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:1015 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
L\!:\!1015 M\!:\!341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1017 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:1017 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
L:1017 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1019\ M:258\ W: Mandatory Feature missing, <221> not found for SEQ ID#:25
L:1019\ M:258\ W: Mandatory Feature missing, <222> not found for SEQ ID#:25
L:1019 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1032 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:26
L:1032 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:26
L:1032 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:1034 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:1042 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:1057 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1061 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1063 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1065 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1067 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1080 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1088 \text{ M}:341 \text{ W}: (46) \text{ "n" or "Xaa" used, for SEQ ID#:28}
L:1090 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1092 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1094 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1096 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1109 \ M:341 \ W: (46) \ "n" \ or "Xaa" \ used, for SEQ ID#:29
L:1111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:1113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
```